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OM nucleic - nucleic search, using sw model

Run on: June 12, 2003, 00:44:04 ; Search time 691 Seconds
(without alignments)
1190.223 Million cell updates/sec

Title: US-09-515-806A-1

Perfect score: 5525

Sequence: 1 tcgcccacgcgtccgcacc.....aatgctttatatactcgca 5525

Scoring table: IDENTITY NUC

Gapop 10.0 , Gapext 1.0

Searched: 870385 seqs, 699768693 residues

Total number of hits satisfying chosen parameters: 1740770

Minimum DB seq length: 0

Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%

Maximum Match 100%

Listing first 45 summaries

Database : Published Applications NA:**

- 1: /cgn2_6/ptodata/1/pubpna/US07_PUBCOMB.seq.*
- 2: /cgn2_6/ptodata/1/pubpna/PCT_NEW_PUB.seq.*
- 3: /cgn2_6/ptodata/1/pubpna/US06_NEW_PUB.seq.*
- 4: /cgn2_6/ptodata/1/pubpna/US06_PUBCOMB.seq.*
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- 13: /cgn2_6/ptodata/1/pubpna/US60_NEW_PUB.seq.*
- 14: /cgn2_6/ptodata/1/pubpna/US60_PUBCOMB.seq.*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Score	Query Match	Length	ID	Description
1	5525	100.0	5525	10	US-09-515-806-1
2	4733	85.7	4989	9	US-09-842-758-3
3	2928.8	53.0	2946	9	US-09-836-392-6
4	2144.6	38.8	2200	10	US-09-925-301-184
5	517.4	9.4	519	10	US-09-988-598-753
6	362.8	6.6	419	9	US-09-918-995-24436
7	248.4	4.5	251	9	US-10-060-036-1707
8	198.6	3.6	19616	10	US-09-764-877-3220
9	191	3.5	147309	10	US-09-742-312-3
10	189.2	3.4	1212	9	US-09-892-877-93
11	189.2	3.4	1212	9	US-09-948-783-92
12	188.2	3.4	326014	10	US-09-731-231A-3
13	187.6	3.4	81001	9	US-09-842-364-1
14	187.6	3.4	81001	10	US-09-751-877-1
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C	22	185.8	3.4	32183	10	US-09-764-869-1494	Sequence 1494, Ap
C	23	185.6	3.4	15297	9	US-10-003-295-3	Sequence 3, Appli
C	24	185.6	3.4	176373	9	US-10-095-407-17	Sequence 17, Appli
C	25	184.8	3.3	25377	9	US-10-061-119-4	Sequence 4, Appli
C	26	183.4	3.3	11172	9	US-10-079-854-231	Sequence 231, App
C	27	183.4	3.3	11172	10	US-09-764-878-231	Sequence 231, App
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C	38	181.6	3.3	416	9	US-09-918-995-16920	Sequence 16920, A
C	39	181.2	3.3	32170	9	US-10-074-095-1108	Sequence 1108, Ap
C	40	181.2	3.3	32170	10	US-09-764-860-1108	Sequence 1108, Ap
C	41	181.2	3.3	44848	9	US-09-988-113-42	Sequence 42, Appli
C	42	181.2	3.3	44848	10	US-09-776-874A-42	Sequence 42, Appli
C	43	180.8	3.3	3422	9	US-09-764-891-6046	Sequence 6046, Ap
C	44	180.8	3.3	3422	9	US-09-764-891-6048	Sequence 6048, Ap
C	45	180.8	3.3	3422	9	US-10-091-438-271	Sequence 271, App

ALIGNMENTS

RESULT 1

US-09-515-806-1

; Sequence 1, Application US/09515806

; Patent No. US20020132321A1

; GENERAL INFORMATION:

; APPLICANT: COOK, WILLIAM J.

; APPLICANT: KAPPELLER-LIBERMANN, ROSANA

; TITLE OF INVENTION: 14790. NOVEL PROTEIN KINASE MOLECULE AND USES THEREFOR

; FILE REFERENCE: 38155-20002.00

; CURRENT APPLICATION NUMBER: US/09/515,806

; CURRENT FILING DATE: 2000-02-29

; NUMBER OF SEQ ID NOS: 32

; SOFTWARE: PatentIn Ver. 2.1

; SEQ ID NO 1

; LENGTH: 5525

; TYPE: DNA

; ORGANISM: Homo sapiens

; FEATURE:

; NAME/KEY: CDS

; LOCATION: (63)..(4991)

US-09-515-806-1

Query Match 100.0%; Score 5525; DB 10; Length 5525;

Best Local Similarity 100.0%; Pred. No. 0;

Matches 5525; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

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Qy	121	CGCAACGACAGGACCAACGAGCTACAGGCGCTCGAGGCCATCTACGCGCGGACTTCCAA	180
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DB 181 TTGTACCTCAAGGCTTAACGTGAAGAATATATGTAAGTGGATTTGAGGTTAAA 240
QY 297 TGCCACCTCACTATCCAGATGATGCTTCTGAATAGAGTTAAATAATGCCAAAGGCTA 356
DB 241 TGCCACCTCACTATCCAGATGATGCTTCTGAATAGAGTTAAATAATGCCAAAGGCTA 300
QY 357 TCAATAGAAAGTGCTCAATTTGTAATAATCTCGCTAGAGAACTGGCCAGAAACACTGT 416
DB 301 TCAATAGAAAGTGCTCAATTTGTAATAATCTCGCTAGAGAACTGGCCAGAAACACTGT 360
QY 417 GGGGAG---GTGATGATCTTTGAACTGGCTTACCACGTGCACTCATTTCTCAGCGAGCAT 473
DB 361 GGGGAGGTAGTGATGATCTTTGAACTGGCTTACCACGTGCACTCATTTCTCAGCGAGCAT 420
QY 474 AACAGCCCCCTCCCAAGTCTTTTCATGAAGAAATGCTGGAAGCGGGCTCAGGAGAG 533
DB 421 AACAGCCCCCTCCCAAGTCTTTTCATGAAGAAATGCTGGAAGCGGGCTCAGGAGAG 480
QY 534 CAGCAGAGCTGTTGAGGCGCAAGCGGAAGAGAGCAGGAGCAACGTGAATCCTGCAT 593
DB 481 CAACAGAGCTGTTGAGGCGCCCAAGCGGAAGAGAGCAGGAGCAACGTGAATCCTGCAT 540
QY 594 GAGATTCAAGAGGAAAGAGAGATAAAGAGAGAAAGAAAGAAATGGCTAAG 653
DB 541 GAGATTCAAGAGGAAAGAGAGATAAAGAGAGAAAGAAAGAAATGGCTAAG 600
QY 654 CAGGAACGTTTGGAAATTCGTAGTTTTCATGAAGAAATGCTGGAAGCGGGCTCAGGAGAG 713
DB 601 CAGGAACGTTTGGAAATTCGTAGTTTTCATGAAGAAATGCTGGAAGCGGGCTCAGGAGAG 660
QY 714 GGAGACACAGAACGGCTGCCATCTACATGGAGCTCTCCTGACTTTGTAGGAAATGGT 773
DB 661 GGAGACACAGAACGGCTGCCATCTACATGGAGCTCTCCTGACTTTGTAGGAAATGGT 720
QY 774 AAACATCGGGCAACTCTCAGGAAGGTCT-----AGGCGAGAACGTCAGTATTCGTGA 827
DB 721 AAACATCGGGCAACTCTCAGGAAGGTCTAGGTTAAGCGAGAACGTCAGTATTCGTGA 780
QY 828 TGTAATAGTAAGATTCCTCGGCTCTTGTGAATTCGTGAATTCGTGAATTCGTGAATTCGTGA 887
DB 781 TGTAATAGTAAGATTCCTCGGCTCTTGTGAATTCGTGAATTCGTGAATTCGTGAATTCGTGA 840
QY 888 GATCAGCTCATGGTGCACAAAGGAAATGATTTGCACTGATGAACACTTGGAAATTA 947
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QY 1008 CAGTGGCAGAAAAAATGGTCTCATCTTACCAGTCAAGAAAAAGAGAAATGATAAG 1067
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QY 1128 CCAATATGATGATGATGATGATGATGATGATGATGATGATGATGATGATGATGATGATGATGAT 1186
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DB 1140 GGACATTTTGTGGAGCAGATTAGTGGGGTCTCTTCTGCTGACACCTGAGGCCACTCAGG 1199
QY 1247 CCCCATCCTGTGATCAGTTCGAGGTGACACAGCTCAGCTCCTGTCAGGCTTTGATTA 1306
DB 1200 CCCCATCCTGTGATCAGTTCGAGGTGACACAGCTCAGCTCCTGTCAGGCTTTGATTA 1259

QY 1307 TCTGCACAGCAATTTCTGTGTGCATGAAGGTCTGAGTGCATCTAATGTCTTGGTGGATGC 1366
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DB 1320 AGAAGGCAACGCTCAAGATTACGGAATATAGCAATTTCTAAGCGCCTCGCAGACATTTGCAAA 1379
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DB 1380 GGAAGGATGTTTGTAGCAAAACCGAGTTCGTTTGTAGCAATGCTCTGCCCTTATAAAAC 1439
QY 1487 GGGGAAAGAGGAGATGTTTGGCGCTTCTGGCCCTTCTGCTGCTGCTCAGCAAGGACA 1546
DB 1440 GGGGAAAGAGGAGATGTTTGGCGCTTCTGGCCCTTCTGCTGCTCAGCAAGGACA 1499
QY 1547 GGAATGTGAGAGTACCTGTGACCAATCCTTAGTGAATTTACCAGCTGACTTTCAAGATTT 1606
DB 1500 GGAATGTGAGAGTACCTGTGACCAATCCTTAGTGAATTTACCAGCTGACTTTCAAGATTT 1559
QY 1607 TCTAAAAGAA--ATGTGTGTGCTTGGATGACAAAGAAAGATGAGTCCCCACAGTGT 1663
DB 1560 TCTAAAAGAAAGATGTTGCTTGGATGACAAAGAAAGATGAGTCCCCACAGTGT 1619
QY 1664 GAAACACAGCTTTATAAATCCCCCAGCCAAAATGCTCTAGTGGAAACAAAGTCTTGAAGA 1723
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QY 1724 TTCTGAGACAAAGATTTATGAGACTGTTATTTCTAGCAACCGGCTACCCAGTGTCTGC 1783
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QY 1784 CTTCTTTTGTAGTACACAGAGACAGTTTCCCGATCTTCTAGTGGAAATTTGAAGAATTTACA 1843
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QY 1904 CTGCTACGAGTGAAGCGCATCCCATCAACCGGCGAGCCGCGAGTTCGGCAGAGATCAA 1963
DB 1857 CTGCTACGAGTGAAGCGCATCCCATCAACCGGCGAGCCGCGAGTTCGGCAGAGATCAA 1916
QY 1964 GGGCGAAGTGAACATCTCTGCTGACGCTGACCATGAGAAACATTTGTCGCTACTCAACGCG 2023
DB 1917 GGGCGAAGTGAACATCTCTGCTGACGCTGACCATGAGAAACATTTGTCGCTACTCAACGCG 1976
QY 2024 CTGGATCGAGCGGACAGCGCGCGGGAGCCGGGAGCCGCGGCTCCCGGCTCCGGGCGC 2083
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DB 2037 CTTGGCCAAAGGATGACCGAGCTGCACGGGGGAGCCGGGAGCGGCTCCCGGCTTGA 2096
QY 2144 CAGCTGAGAGGCGCGCGCGGCTCCCATCTCAGCAGCTCGGTGGAGTGGAGCACTTC 2203
DB 2097 CAGCTGAGAGGCGCGCGCGGCTCCCATCTCAGCAGCTCGGTGGAGTGGAGCACTTC 2156
QY 2204 GGGCGAGCGCTCGGCGAGTCCCGGCTTCCCGGCTCCCGGCTCCCGGCTCCCGGCTCCCGGCT 2263
DB 2157 GGGCGAGCGCTCGGCGAGTCCCGGCTTCCCGGCTCCCGGCTCCCGGCTCCCGGCTCCCGGCT 2216
QY 2264 GGAAGCAGAGGAGGAGCAGCGGTGGGCTCTTCTCCAGTCTCTTCTGCTGCTTCTGCTTCTGCT 2323
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QY 2324 TTCTGAAAGTGTATTTATCTTTGACAAATGAAAGTGAAGAAACAGTAAAGTCAAGATCAGGA 2383
DB 2277 TTCTGAAAGTGTATTTATCTTTGACAAATGAAAGTGAAGAAACAGTAAAGTCAAGATCAGGA 2336
QY 2384 TGAAGATTGCAATGAAAGAAATGGCTGCCATGAAAGTGAAGCCATCAGTGCAGCTGAGGC 2443

QY 4583 TGCAGTCAAAATCTGAAGGGGTCAATTTCTAAATGCTTCAGGTTTGTGTTGAAATCCATGG 4642
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 QY 4557 TGCAGTCAAAATCTGAAGGGGTCAATTTCTAAATGCTTCAGGTTTGTGTTGAAATCCATGG 4616
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 QY 4643 AGCAACAGTGGTTCATTTGAGTGTGCTAGCCCGGAGAAAGTGTGACGACACTAG 4702
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 QY 4940 CAAAGTAGAAAAAGGTGCTGTGCTATTTCTGTACAGCTATAGAGATGACTACTACAG 4999
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 QY 4917 CAAAGTAGAAAAAGGTGCTGTGCTATTTCTGTACAGCTATAGAGATGACTACTACAG 4976
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 QY 5000 AATCTTATTTTAA 5012
 Db |||||
 QY 4977 AATCTTATTTTAA 4989

RESULT 3

US-09-836-392-6
 ; Sequence 6, Application US/09836392
 ; Patent No. US20020173458A1
 ; GENERAL INFORMATION:
 ; APPLICANT: Ruben et al.
 ; TITLE OF INVENTION: Protein Tyrosine Kinase Receptor Polynucleotides, Polypeptides, a
 ; TITLE OF INVENTION: Antibodies
 ; FILE REFERENCE: PT020P1
 ; CURRENT APPLICATION NUMBER: US/09/836,392
 ; CURRENT FILING DATE: 2001-04-18
 ; PRIOR APPLICATION NUMBER: PCT/US00/28066
 ; PRIOR FILING DATE: 2000-10-11
 ; PRIOR APPLICATION NUMBER: 60/159,542
 ; PRIOR FILING DATE: 1999-10-15
 ; PRIOR APPLICATION NUMBER: 60/165,914
 ; PRIOR FILING DATE: 1999-11-17
 ; PRIOR APPLICATION NUMBER: 60/189,027
 ; PRIOR FILING DATE: 2000-03-14
 ; NUMBER OF SEQ ID NOS: 34
 ; SOFTWARE: PatentIn Ver. 2.0
 ; SEQ ID NO 6
 ; LENGTH: 2946
 ; TYPE: DNA
 ; ORGANISM: Homo sapiens
 US-09-836-392-6

Query Match 53.0%; Score 2928.8; DB 9; Length 2946;
 Best Local Similarity 99.9%; Pred. No. 0;
 Matches 2930; Conservative 0; Mismatches 2; Indels 0; Gaps 0;

QY 2485 GCACCTTTACGAGACACCAATGACAGGACTGTATCGAGACACCGTCAGACTCTGGAGC 2544
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 QY 12 GCACCTTTACGAGACACCAATGACAGGACTGTATCGAGACACCGTCAGACTCTGGAGC 71
 Db |||||
 QY 2545 TTTTTCAGAGATCTGGATGATAGCTTATATCCATGAGAAGGAATGATTCACCGGG 2604
 Db |||||
 QY 72 TTTTTCAGAGATCTGGATGATAGCTTATATCCATGAGAAGGAATGATTCACCGGG 131
 Db |||||

QY 2605 ATTTGAAGCCTCTCAACATTTTGGATTTCTGATGACCATGTGAAATAGGTGATTTTG 2664
 Db |||||
 QY 132 ATTTGAAGCCTCTCAACATTTTGGATTTCTGATGACCATGTGAAATAGGTGATTTTG 191
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 QY 2665 GTTTGGCGACAGACCATCTAGCCTTTCTGCTGACAGCAAAACAGACGATCAGACAGGAG 2724
 Db |||||
 QY 192 GTTTGGCGACAGACCATCTAGCCTTTCTGCTGACAGCAAAACAGACGATCAGACAGGAG 251
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 QY 2725 ACTTGATTAAGTCAGACCCCTTCAGGTCACTTAATCTGGATGTTGGCACTGCTCTCTATG 2784
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 QY 252 ACTTGATTAAGTCAGACCCCTTCAGGTCACTTAATCTGGATGTTGGCACTGCTCTCTATG 311
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 QY 2785 TAAGCCAGAGTCCAAAGGAAGCACCAATCTGCATACAACCCAGAAAGTGGATCTCTTCA 2844
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 QY 312 TAAGCCAGAGTCCAAAGGAAGCACCAATCTGCATACAACCCAGAAAGTGGATCTCTTCA 371
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 QY 2845 GCCTGGGAATTAATCTTCTTTGAGATGCTCTATCACCCCATGGTTCACGGCTTCAGAAAAGA 2904
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 QY 372 GCCTGGGAATTAATCTTCTTTGAGATGCTCTATCACCCCATGGTTCACGGCTTCAGAAAAGA 431
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 QY 2905 TCTTTTGTCTCAACCAACTCAGAGATCCACTTTCGCTTAAGTTTCCAGAAAGTCTTGACG 2964
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 QY 2965 ATGGAGAGCATGCCAAAGCAGAAATCAGTCATCTCTGCTGTTGAACCAAGTCCAGAA 3024
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 QY 492 ATGGAGAGCATGCCAAAGCAGAAATCAGTCATCTCTGCTGTTGAACCAAGTCCAGAA 551
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 QY 3025 AACGGCCACAGCCACAGAACTGCTCAAGAGTGAGTCTGCTCCCAACCCAGATGGAGG 3084
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 QY 3385 GCGGATGCTGATGCTTCTTTGACCTGCGGATCCCTTTTGAAGATATGTGGCAA 3444
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 QY 912 GCGGATGCTGATGCTTCTTTGACCTGCGGATCCCTTTTGAAGATATGTGGCAA 971
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 QY 3505 TAGATCGATTTTATCCCAAGAACTTCTGGAGTGTGATTTGATTTGTCTACCTTCTTACCA 3564
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 QY 1032 TAGATCGATTTTATCCCAAGAACTTCTGGAGTGTGATTTGATTTGTCTACCTTCTTACCA 1091
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 QY 3565 CCAACAGCTTCTGCGCACTGCTGAAATTTATCTACCTATCTATGAAATCATCCAGAGT 3624
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 QY 1092 CCAACAGCTTCTGCGCACTGCTGAAATTTATCTACCTATCTATGAAATCATCCAGAGT 1151
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 QY 3625 TTCCAGCACTTCAGAAAGAAATTTACAGTATTTATTTTGAACCACTACCATCTGTTTGAAG 3684
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 QY 1152 TTCCAGCACTTCAGAAAGAAATTTACAGTATTTATTTTGAACCACTACCATCTGTTTGAAG 1211
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 QY 3685 CAATACCTTTACACTGTGGATCCCAAGAGATAAACTCAGTCAAGTCTACATTTCTGT 3744
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; NAME/KEY: misc_feature
; LOCATION: (2181)
; OTHER INFORMATION: n equals a,t,g, or c
; NAME/KEY: misc_feature
; LOCATION: (2184)
; OTHER INFORMATION: n equals a,t,g, or c
US-09-925-301-184

Query Match      38.8%; Score 2144.6; DB 10; Length 2200;
Best Local Similarity 99.0%; Pred. No. 0;
Matches 2173; Conservative 2; Mismatches 16; Indels 3; Gaps 2;

QY 3206 CAGGACATCTGAAGGCACTTCTCAATCCGTACAGCCAAAGATGACAGCAGATGTGTG 3265
Db 9 CAGGACATCTGAAGGCACTTCTCAATCCGTACAGCCAAAGATGACAGCAGATGTGTG 68

QY 3266 TGAACCATCATCCGCACTTTTAAAGACATGAGTGTTCAGTGTGTACTCCACTACT 3325
Db 69 TGAACCATCATCCGCACTTTTAAAGACATGAGTGTTCAGTGTGTACTCCACTACT 128

QY 3326 GCTTCCCGAAACAGACAATATATGAGCAACAAGAGCTGCCCTATTTCATGGACCACAG 3385
Db 129 GCTTCCCGAAACAGACAATATATGAGCAACAAGAGCTGCCCTATTTCATGGACCACAG 188

QY 3386 CGGATGCTGTGTGATGCTTCTCTTTTGACCTCGCGATCCCTTTTGCAGATATGTGGCAAG 3445
Db 189 CGGATGCTGTGTGATGCTTCTCTTTTGACCTCGCGATCCCTTTTGCAGATATGTGGCAAG 248

QY 3446 AAATAATATATTTAAACGATATCTCATAGAACGTGTTCAGGCCGCGCAAGTT 3505
Db 249 AAATAATATATTTAAACGATATCTCATAGAACGTGTTCAGGCCGCGCAAGTT 308

QY 3506 AGATCGATTTTATCCCAAGAACTTCTGGAGTGTGCAATTTGATATTTGCTTCTTACCAC 3565
Db 309 AGATCGATTTTATCCCAAGAACTTCTGGAGTGTGCAATTTGATATTTGCTTCTTACCAC 368

QY 3566 CAACAGCTTTCTGCGCACTCTGAAATATCTACATCTATCTATGAATCATCCAGAGTT 3625
Db 369 CAACAGCTTTCTGCGCACTCTGAAATATCTACATCTATCTATGAATCATCCAGAGTT 428

QY 3626 TCCAGCACTTTCAGAAAGAAATACAGTATTTATTTGAACCATACCATGTTATTGAAGC 3685
Db 429 TCCAGCACTTTCAGAAAGAAATACAGTATTTATTTGAACCATACCATGTTATTGAAGC 488

QY 3686 AATACCTTTACACTGTGGATCCGAGAGATAAATCAAGTCAAGTCTACATTTCTGTA 3745
Db 489 AATACCTTTACACTGTGGATCCGAGAGATAAATCAAGTCAAGTCTACATTTCTGTA 548

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Db 549 TGATGCTGTGACAGAGAGCTGACGAGGAGAGAGTGGAGCTTAAATTTTGAATCTGTC 608

QY 3806 TTTGCTTTCTAATAGTCTGTGCTGACTCTACAGTATTTATTTGAACAGAGGAGATTGCA 3865
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QY 3926 GAAATATGGCTTAAAGACCTAGAGAGGTTGTGTGACGTTTGAAGAACTCGGCATCAA 3985
Db 729 GAAATATGGCTTAAAGACCTAGAGAGGTTGTGTGACGTTTGAAGAACTCGGCATCAA 788

QY 3986 GTTACAGGCTTTGATCAATTTGGGCTTGGTTTACAGGTGACGACGACCAATGGAAATCAT 4045
Db 789 GTTACAGGCTTTGATCAATTTGGGCTTGGTTTACAGGTGACGACGACCAATGGAAATCAT 848

QY 4046 CTTCCAGTTTGGCTTTTTCATCAACAGAGGCAAGGGCTGTACCTGAAATCCTCGAGC 4105
Db 849 CTTCCAGTTTGGCTTTTTCATCAACAGAGGCAAGGGCTGTACCTGAAATCCTCGAGC 908

QY 4106 TGGAGGACAGATATGACCTGCTGATTTCCCCAGTTTATAGAGGCCACAGCTCTGGGCCAGT 4165
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909 TGGAGGACAGATATGACCTGCTGATTTCCCCAGTTTATAGAGGCCACAGCTCTGGGCCAGT 968
4166 TCCCACTGCCATTGGGGTTCAGCATAGCTATAGACAAGATATCTGCTGTCTCTCAACAT 4225
969 TCCCACTGCCATTGGGGTTCAGCATAGCTATAGACAAGATATCTGCTGTCTCTCAACAT 1028
4226 GGAGGAATCTGTACAAATAGCTCTGTGACCTCTCTGTTTAAAGTGTGTGTCAGATGTC 4285
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4466 GAAAGAAAGGACAGACAGAAAGCGTGTCTGGAGTCTGAACTTTGTGGACCACTGACTGCA 4525
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1689 ACGCTTCCCAAGCAAGATACCTCAATTTAGTCTGTGATGAAATTTTAAACATCAAGT 1748
4946 AGAAAAAGAGTGTGCTGTGCTTATTTCTGTACAGCTATAGAGATGACTACTACAGAACTT 5005
1749 AGAAAAAGAGTGTGCTGTGCTTATTTCTGTACAGCTATAGAGATGACTACTACAGAACTT 1808
5006 ATTTTAAACCTTAAAGAACTGTGTTAACTTCAATTTAACTTAACTTAACTTAACTTAACT 5065
1809 ATTTTAAACCTTAAAGAACTGTGTTAACTTCAATTTAACTTAACTTAACTTAACTTAACT 1868
5066 TAATGGAATGTGTGATCAATTTTAACTTAACTTAACTTAACTTAACTTAACTTAACTTAACT 5125
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5126 AGTGGCTCACACCTTTAATCCAGCACTTTGGGAAGCCAGGAGGAGGAGGAGGAGGAGGAGG 5185
1929 AGTGGCTCACACCTTTAATCCAGCACTTTGGGAAGCCAGGAGGAGGAGGAGGAGGAGGAGG 1988
5186 CCAGGAGTTTGGAGCAGGCTGTGAGCAACAAAGCAAGCCCCATCTCTATATAAACTTAACT 5245
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APPLICANT: Jiang, Yufu
; TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY
; TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY
; FILE REFERENCE: 210121.566
; CURRENT APPLICATION NUMBER: US/10/060,036
; CURRENT FILING DATE: 2002-01-30
; NUMBER OF SEQ ID NOS: 4560
; SOFTWARE: FASTSEQ for Windows Version 4.0
; SEQ ID NO 1707
; LENGTH: 251
; TYPE: DNA
; ORGANISM: Homo sapiens
; FEATURE:
; NAME/KEY: misc_feature
; LOCATION: 244
; OTHER INFORMATION: n = A,T,C or G
US-10-060-036-1707

Query Match 4.5%; Score 248.4; DB 9; Length 251;
Best Local Similarity 99.2%; Pred. No. 4.6e-57;
Matches 249; Conservative 0; Mismatches 2; Indels 0; Gaps 0;

QY 4724 ACAACTCGACTTCAGACCTCCCTGCGCAACTTACATCAGAAAGCAGTGAATGAAAT 4783
Db 251 ACAACTCGACTTCAGACCTCCCTGCGCAACTTACATCAGAAAGCAGTGAATGAAAT 192
QY 4784 TCTGCTGTGGATCTTACCCAAAGAAACAATATTACAGTTTATCATTTAGAGTGGATGC 4843
Db 191 TCTGCTGTGGATCTTACCCAAAGAAACAATATTACAGTTTATCATTTAGAGTGGATGC 132
QY 4844 TGATGAACAGGCAATTAACAACATGTGAAGCAGCTGCTGCACGCCCTGCCAAAGCAAAG 4903
Db 131 TGATGAACAGGCAATTAACAACATGTGAAGCAGCTGCTGCACGCCCTGCCAAAGCAAAG 72
QY 4904 ATACCTCAAAATAGCTCTGTGATGAATTTATTAACATCAAGTAGAAAAAGGTGTCTGT 4963
Db 71 ATACCTCAAAATAGCTCTGTGATGAATTTATTAACATCAAGTAGAAAAAGGTGTCTGT 12
QY 4964 GCTATTCTGT 4974
Db 11 GCTATTCTGT 1

RESULT 8
US-09-764-877-3220/c
; Sequence 3220, Application US/09764877
; Patent No. US20020147140A1
; GENERAL INFORMATION:
; APPLICANT: Rosen et al.
; TITLE OF INVENTION: Nucleic Acids, Proteins, and Antibodies
; FILE REFERENCE: PC005
; CURRENT APPLICATION NUMBER: US/09/764,877
; CURRENT FILING DATE: 2001-01-17
; Prior application data removed - refer to PALM or file wrapper
; NUMBER OF SEQ ID NOS: 4031
; SOFTWARE: Patent in Ver. 2.0
; SEQ ID NO 3220
; LENGTH: 19616
; TYPE: DNA
; ORGANISM: Homo sapiens
US-09-764-877-3220

Query Match 3.6%; Score 198.6; DB 10; Length 19616;
Best Local Similarity 82.4%; Pred. No. 3.8e-42;
Matches 253; Conservative 0; Mismatches 49; Indels 5; Gaps 2;

QY 5114 GAGCTGGTTCAGTGGCTTACACCTTTATCCAGCAGCTTTGGGAAGCAAGCGAGAA 5173
Db 18954 GAGCTGGTTCAGTGGCTTACACCTTTATCCAGCAGCTTTGGGAAGCAAGCGAGAA 18295
QY 5174 GACTGCTTGAACACAGGAGTTTGAGACCGCTGAGCAACAAAGCAGACCCCTCTCTA 5233
Db 18294 GACGCTTGACCCAGGGGTTTGAGACCGCTGAGCAGCATGGCAAAACCCCTCTCTA 18235

QY 5234 TAAAACTAAAAAATAGTTGGGCATGCTGGCACAATGCTAGTCCAGCTACTCCAG 5293
Db 18234 CAAAAAATCAAAAAATAGTTGGGTGTGGCTATAGTCCAGCTACTCCAG 18175
QY 5294 AGGCTGAGAT---GGATCATCTGAGCCTCAGAGGCTTGGAGCTGAGCTGTGACT 5349
Db 18174 AGGCTGAGGATGAGGATCACTGAGCCTGGGAGGTAGAGGCTGAGTGACCGCAT 18115
QY 5350 GCGCCACTGCTCAGTCTGGGACAACAGACCAAGACCTCTCTTAAAAAAGAA 5409
Db 18114 GCACCACTGCTCAGCTCCAGCCTGGG-CAACAGAGCAAGACCTCTCTGAAAAAAGAA 18056
QY 5410 AAAAAA 5416
Db 18055 AAAAAA 18049

RESULT 9
US-09-742-312-3
; Sequence 3, Application US/09742312
; Patent No. US20020045166A1
; GENERAL INFORMATION:
; APPLICANT: CHANDRAMOULISARAN, Ishwar et al
; TITLE OF INVENTION: ISOLATED HUMAN TRANSPORTER PROTEINS,
; TITLE OF INVENTION: NUCLEIC ACID MOLECULES ENCODING HUMAN TRANSPORTER PROTEINS,
; TITLE OF INVENTION: AND USES THEREOF
; FILE REFERENCE: CL000838
; CURRENT APPLICATION NUMBER: US/09/742,312
; CURRENT FILING DATE: 2000-12-22
; NUMBER OF SEQ ID NOS: 4
; SOFTWARE: FastSEQ for Windows Version 4.0
; SEQ ID NO 3
; LENGTH: 147309
; TYPE: DNA
; ORGANISM: Human
; FEATURE:
; NAME/KEY: misc_feature
; LOCATION: (1)-(147309)
; OTHER INFORMATION: n = A,T,C or G
US-09-742-312-3

Query Match 3.5%; Score 191; DB 10; Length 147309;
Best Local Similarity 66.8%; Pred. No. 1.9e-39;
Matches 272; Conservative 0; Mismatches 135; Indels 0; Gaps 0;

QY 5076 TTGTACATTCATCAATAATTTAAATTTAAATTTCTAAGAGAGCTGGTGCAGTGCAC 5135
Db 57224 TTGTAAAAAATTTAAATTTAAATTTAAATTTCTAAGAGAGCTGGTGCAGTGCAC 57283
QY 5136 ACCTTTAATCCAGCAGCTTTGGGAGCCAGCAGGAGAGCTGCTTTGAAACGAGGATTT 5195
Db 57284 GCCTATAATCCAGCAGCTTTGGGAGCCAGCAGGAGAGCTGCTTTGAAACGAGGATTT 57343
QY 5196 GAGACGAGCTGAGCAACAAAGACCCATCTCTATAAAAACTTAAAAATTTAGTTG 5255
Db 57344 GAGACGAGCTGAGCAACAAAGACCCATCTCTATAAAAACTTAAAAATTTAGTTG 57403
QY 5256 GGCATGGTGGCAGATGCTGTAGTCCAGCTACTCCAGGCTGAGATGGATCTCAG 5315
Db 57404 GGCATGGTGGTGTGTGCTGTAGTCCAGCTACTCCAGGCTGAGGAGGAGATTTG 57463
QY 5316 CCTCAGGAGGTTGAGGCTGAGCTGAGCTGAGCTGCGCCACTGCACTCCAGTCTGGGACA 5375
Db 57464 ACCAAGAGGAGGAGGTTGAGCTGAGCTGAGCTGAGCTGAGCTGAGCTGAGCTGAG 57523
QY 5376 ACAGAGCAAGACCTCTTTAAAAAAGAAAAAATTTTCTTAAAGAGCTG 5435
Db 57524 CAAGAGTGAATCTCCATCTCAGAAAAAATTTTCTTAAAGAGCTG 57583
QY 5436 TCCTACAAAGTTGAGCTTTGTTAGTTTCTATGTGTAATATATATA 5482
Db 57584 TAATCAATTTCTTTTGGGTGGAGCAGCAATTTGTTCTTTTAAA 57630

RESULT 10

US-09-892-877-93

; Sequence 93, Application US/09892877
; Publication No. US20030077809A1

; GENERAL INFORMATION:

; APPLICANT: Ruben et. al.

; TITLE OF INVENTION: 97 Human secreted proteins

; FILE REFERENCE: P2028P1

; CURRENT APPLICATION NUMBER: US/09/892,877

; CURRENT FILING DATE: 2001-06-28

; PRIOR APPLICATION NUMBER: EARLIER APPLICATION NUMBER: US/09/437,658

; PRIOR FILING DATE: EARLIER FILING DATE: 1999-11-10

; NUMBER OF SEQ ID NOS: 461

; SOFTWARE: PatentIn Ver. 2.0

; SEQ ID NO 93

; TYPE: DNA

; ORGANISM: Homo sapiens

; US-09-892-877-93

Query Match 3.4%; Score 189.2; DB 9; Length 1212;

Best Local Similarity 76.7%; Pred. No. 1.8e-40;

Matches 247; Conservative 7; Mismatches 65; Indels 3; Gaps 2;

Qy	5095	TAAATTAATTTCTAAGAGAGGCTGGGTGCAGTGGCTCACACCTTTAATCCAGCAGCTT	5154
Db	892	TACATATGAATAATAGGTACAGCCAGGTGTAGTGGCTCACACCTGTATATCCAGCAGCTT	951
Qy	5155	TGGGAAGCCCAAGCAGGAGACTGCTTTGAAACCCAGGAGTTTGAGACCAGCCTTGAGCAACA	5214
Db	952	GGGGAAGCCGARGTGGGTGATTTGTTGAGCCAGGAGTTTGAGACCAGCCTTGAGCAACA	1011
Qy	5215	AAGCAAGACCCCATCTCTATAAACTATAAACTATAAACTATAAACTATAAACTATAAACT	5274
Db	1012	TGGTGAACCCCATCTCTACAAAAATMCAAAAAATMCAAAAAATMCAAAAAATMCAAAAAAT	1071
Qy	5275	GTAGTCCAGCTACTCCAGAGGCTGAGATGG--ATCATCTGAGCCTCAGAGGTTTGAGGC	5332
Db	1072	GTAGTCTCAGCTACTCAGAGGCTGAGATGG--ATCATCTGAGCCTCAGAGGTTTGAGGC	5332
Qy	5333	TGCAGTGAGCTGTGACTTGGCCACTGCACCTCCAGTCTGGGACAAACAGAGCAAGCCCTGT	5392
Db	1132	TGCAGTGAGCTGTGATCCTGCCACTGSACTCCAGCCTGGG--TGACAGGCGCAAGACCCCTGT	1190
Qy	5393	CTTAAAAAAGAAAAA 5414	
Db	1191	YTWAAAAAAGAAAAA 1212	

RESULT 11

US-09-948-783-92

; Sequence 92, Application US/09948783

; Publication No. US20030100051A1

; GENERAL INFORMATION:

; APPLICANT: Ruben et. al.

; TITLE OF INVENTION: 97 Human secreted proteins

; FILE REFERENCE: P2028P2

; CURRENT APPLICATION NUMBER: US/09/948,783

; CURRENT FILING DATE: 2001-09-10

; PRIOR APPLICATION NUMBER: 60/231,846

; PRIOR FILING DATE: 2000-09-11

; PRIOR APPLICATION NUMBER: 09/892,877

; PRIOR FILING DATE: 2001-06-28

; PRIOR APPLICATION NUMBER: 09/437,658

; PRIOR FILING DATE: 1999-11-10

; PRIOR APPLICATION NUMBER: PCT/US99/09847

; PRIOR FILING DATE: 1999-05-06

; PRIOR APPLICATION NUMBER: 60/085,093

; PRIOR FILING DATE: 1998-05-12

; PRIOR APPLICATION NUMBER: 60/085,094

; PRIOR FILING DATE: 1998-05-12

; PRIOR APPLICATION NUMBER: 60/085,105

; PRIOR FILING DATE: 1998-05-12

; PRIOR APPLICATION NUMBER: 60/085,180

; PRIOR FILING DATE: 1998-05-12

; PRIOR APPLICATION NUMBER: 60/085,927

; PRIOR FILING DATE: 1998-05-18

; PRIOR APPLICATION NUMBER: 60/085,906

; PRIOR FILING DATE: 1998-05-18

; PRIOR APPLICATION NUMBER: 60/085,924

; PRIOR FILING DATE: 1998-05-18

; PRIOR APPLICATION NUMBER: 60/085,922

; PRIOR FILING DATE: 1998-05-18

; PRIOR APPLICATION NUMBER: 60/085,921

; PRIOR FILING DATE: 1998-05-18

; PRIOR APPLICATION NUMBER: 60/085,923

; PRIOR FILING DATE: 1998-05-18

; PRIOR APPLICATION NUMBER: 60/085,925

; PRIOR FILING DATE: 1998-05-18

; PRIOR APPLICATION NUMBER: 60/085,928

; PRIOR FILING DATE: 1998-05-18

; PRIOR APPLICATION NUMBER: 60/085,920

; PRIOR FILING DATE: 1998-05-18

; NUMBER OF SEQ ID NOS: 465

; SOFTWARE: PatentIn Ver. 2.0

; SEQ ID NO 92

; LENGTH: 1212

; TYPE: DNA

; ORGANISM: Homo sapiens

; US-09-948-783-92

Query Match

Best Local Similarity 76.7%; Pred. No. 1.8e-40;

Matches 247; Conservative 7; Mismatches 65; Indels 3; Gaps 2;

Qy	5095	TAAATTAATTTCTAAGAGAGGCTGGGTGCAGTGGCTCACACCTTTAATCCAGCAGCTT	5154
Db	892	TACATATGAATAATAGGTACAGCCAGGTGTAGTGGCTCACACCTGTATATCCAGCAGCTT	951
Qy	5155	TGGGAAGCCCAAGCAGGAGACTGCTTTGAAACCCAGGAGTTTGAGACCAGCCTTGAGCAACA	5214
Db	952	GGGGAAGCCGARGTGGGTGATTTGTTGAGCCAGGAGTTTGAGACCAGCCTTGAGCAACA	1011
Qy	5215	AAGCAAGACCCCATCTCTATAAACTATAAACTATAAACTATAAACTATAAACTATAAACT	5274
Db	1012	TGGTGAACCCCATCTCTACAAAAATMCAAAAAATMCAAAAAATMCAAAAAATMCAAAAAAT	1071
Qy	5275	GTAGTCCAGCTACTCCAGAGGCTGAGATGG--ATCATCTGAGCCTCAGAGGTTTGAGGC	5332
Db	1072	GTAGTCTCAGCTACTCAGAGGCTGAGATGG--ATCATCTGAGCCTCAGAGGTTTGAGGC	5332
Qy	5333	TGCAGTGAGCTGTGACTTGGCCACTGCACCTCCAGTCTGGGACAAACAGAGCAAGCCCTGT	5392
Db	1132	TGCAGTGAGCTGTGATCCTGCCACTGSACTCCAGCCTGGG--TGACAGGCGCAAGACCCCTGT	1190
Qy	5393	CTTAAAAAAGAAAAA 5414	
Db	1191	YTWAAAAAAGAAAAA 1212	

RESULT 12

US-09-731-231A-3/c

; Sequence 3, Application US/09731231A

; Patent No. US20020082189A1

; GENERAL INFORMATION:

; APPLICANT: GUEGLER, Karl et al

; TITLE OF INVENTION: ISOLATED HUMAN KINASE PROTEINS, NUCLEIC

; TITLE OF INVENTION: ACID MOLECULES ENCODING HUMAN KINASE PROTEINS, AND USES

; TITLE OF INVENTION: THEREOF

; FILE REFERENCE: CL001007

; CURRENT APPLICATION NUMBER: US/09/731,231A

; CURRENT FILING DATE: 2000-12-07

; NUMBER OF SEQ ID NOS: 6

; SOFTWARE: FastSeq for Windows Version 4.0

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; SEQ ID NO 3
; LENGTH: 326014
; TYPE: DNA
; ORGANISM: Human
; FEATURE:
; NAME/KEY: misc feature
; LOCATION: (1)...(326014)
; OTHER INFORMATION: n = A,T,C or G
US-09-731-231A-3

Query Match      3.4%; Score 188.2; DB 10; Length 326014;
Best Local Similarity 77.8%; Pred. No. 28-38;
Matches 253; Conservative 0; Mismatches 68; Indels 4; Gaps 2;

QY 5101 TAAATTCCTAAGAGAGGCTGGTGCAGTGGCTGCACACCTTTAATCCAGCAGCTTTGGGAA 5160
Db      |||||
QY 36647 TAAAGCTCCATTCAGCTAGATGAGTGGCTCACAACGTGTAATCTAGCAGCTTTGTGAG 36588
Db      |||||
QY 5161 GCCAAGGAGGAGAGTCTGTTGAAACCCAGGAGTTTGAGACCCAGCTGAGCAACAAAGCAA 5220
Db      |||||
QY 36587 GCCAAGGAGGAGAGTCTGTTGAAACCCAGGAGTTTGAGACCCAGCTGAGCAACAAAGCAA 36528
Db      |||||
QY 5221 GACCCATCTCTATAAACTAAATAAATTTAGTGGCATGTGGCATGCGCTGTAGTC 5280
Db      |||||
QY 36527 GACCCCTCTCTATAAAAGTAAATAAATTTAGTGGCATGTGGCATGCGCTGTAGTC 36468
Db      |||||
QY 5281 CCAGCTACTCCAGAGCTGAGATGATCATCTG---AGCCTCAGGAGGTTGAGGCTGCAG 5337
Db      |||||
QY 36467 CCAGCTACTTGGAGGCTGAGCAGGAGGACTGCCTAACCTTAGAGTTTGAGGCTGCAG 36408
Db      |||||
QY 5338 TGAGCTGTGACTGCGCAGCTGCACTGCTGAGTGGGAGCAACAGAGCAAGCCCTGCTTAA 5397
Db      |||||
QY 36407 TGAGCAGTGTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCTGCT 36349
Db      |||||
QY 5398 AAAAAAAGAAAAAATTTTTT 5422
Db      |||||
QY 36348 AAAAAAAGAAAAAATTTTTT 36324
Db      |||||

RESULT 13
US-09-842-364-1/c
; Sequence 1, Application US/09842364
; Publication No. US20030032783A1
; GENERAL INFORMATION:
; APPLICANT: Yen-Potin, Frances
; APPLICANT: Denison, Blake
; APPLICANT: Bout, Barbara
; APPLICANT: Bihain, Bernard
; APPLICANT: Dumas Milne Edwards, Jean-Baptiste
; APPLICANT: Duclert, Aymeric
; APPLICANT: Bougueret, Lydie
; TITLE OF INVENTION: APOLOPROTEIN A-IV-RELATED PROTEIN: POLYPEPTIDE, POLYNUCLEOTIDE
; FILE REFERENCE: GENSET 50CP2C
; CURRENT APPLICATION NUMBER: US/09/842,364
; CURRENT FILING DATE: 2001-04-25
; PRIOR APPLICATION NUMBER: US 09/599,362
; PRIOR FILING DATE: 2000-06-21
; PRIOR APPLICATION NUMBER: PCT/IB99/02058
; PRIOR FILING DATE: 1998-12-20
; PRIOR APPLICATION NUMBER: US 09/469,099
; PRIOR FILING DATE: 1998-12-21
; PRIOR APPLICATION NUMBER: US 60/113,686
; PRIOR FILING DATE: 1998-12-22
; PRIOR APPLICATION NUMBER: US 60/141,032
; PRIOR FILING DATE: 1998-06-25
; NUMBER OF SEQ ID NOS: 6
; SOFTWARE: Patent.pm
; SEQ ID NO 1
; LENGTH: 81001
; TYPE: DNA
; ORGANISM: Homo sapiens
; FEATURE:
; NAME/KEY: misc feature
; LOCATION: 10946..12946
; OTHER INFORMATION: 5'regulatory region
; NAME/KEY: exon
; LOCATION: 12947..12958
; OTHER INFORMATION: exon 1
; NAME/KEY: exon
; LOCATION: 13470..13526
; OTHER INFORMATION: exon 2
; NAME/KEY: exon
; LOCATION: 13641..13752
; OTHER INFORMATION: exon 3
; NAME/KEY: exon
; LOCATION: 14271..15968
; OTHER INFORMATION: exon 4
; NAME/KEY: misc feature
; LOCATION: 15969..17969
; OTHER INFORMATION: 3'regulatory region
; NAME/KEY: allele
; LOCATION: 1239
; OTHER INFORMATION: 20-828-311 : polymorphic base C or T
; NAME/KEY: allele
; LOCATION: 12347
; OTHER INFORMATION: 17-42-319 : polymorphic base C or T
; NAME/KEY: allele
; LOCATION: 15241
; OTHER INFORMATION: 17-41-250 : polymorphic base C or T
; NAME/KEY: allele
; LOCATION: 42218
; OTHER INFORMATION: 20-841-149 : polymorphic base A or G
; NAME/KEY: allele
; LOCATION: 45442
; OTHER INFORMATION: 20-842-115 : polymorphic base A or G
; NAME/KEY: allele
; LOCATION: 77058
; OTHER INFORMATION: 20-853-415 : polymorphic base C or T
; NAME/KEY: primer_bind
; LOCATION: 929..949
; OTHER INFORMATION: 20-828.pu
; NAME/KEY: primer_bind
; LOCATION: 1357..1377
; OTHER INFORMATION: 20-828.rp complement
; NAME/KEY: primer_bind
; LOCATION: 12029..12050
; OTHER INFORMATION: 17-42.pu
; NAME/KEY: primer_bind
; LOCATION: 12581..12603
; OTHER INFORMATION: 17-42.rp complement
; NAME/KEY: primer_bind
; LOCATION: 14992..15012
; OTHER INFORMATION: 17-41.pu
; NAME/KEY: primer_bind
; LOCATION: 15460..15482
; OTHER INFORMATION: 17-41.rp complement
; NAME/KEY: primer_bind
; LOCATION: 42070..42090
; OTHER INFORMATION: 20-841.pu
; NAME/KEY: primer_bind
; LOCATION: 42572..42591
; OTHER INFORMATION: 20-841.rp complement
; NAME/KEY: primer_bind
; LOCATION: 45328..45347
; OTHER INFORMATION: 20-842.pu
; NAME/KEY: primer_bind
; LOCATION: 45863..45883
; OTHER INFORMATION: 20-842.rp complement
; NAME/KEY: primer_bind
; LOCATION: 76644..76664
; OTHER INFORMATION: 20-853.pu
; NAME/KEY: primer_bind
; LOCATION: 77166..77185
; OTHER INFORMATION: 20-853.rp complement
; NAME/KEY: primer_bind
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; NAME/KEY: allele
; LOCATION: 77058
; OTHER INFORMATION: 20-853-415 : polymorphic base C or T
; NAME/KEY: primer_bind
; LOCATION: 929..949
; OTHER INFORMATION: 20-828.pu
; NAME/KEY: primer_bind
; LOCATION: 1357..1377
; OTHER INFORMATION: 20-828.rp complement
; NAME/KEY: primer_bind
; LOCATION: 12029..12050
; OTHER INFORMATION: 17-42.pu
; NAME/KEY: primer_bind
; LOCATION: 12581..12603
; OTHER INFORMATION: 17-42.rp complement
; NAME/KEY: primer_bind
; LOCATION: 14992..15012
; OTHER INFORMATION: 17-41.pu
; NAME/KEY: primer_bind
; LOCATION: 15460..15482
; OTHER INFORMATION: 17-41.rp complement
; NAME/KEY: primer_bind
; LOCATION: 42070..42090
; OTHER INFORMATION: 20-841.pu
; NAME/KEY: primer_bind
; LOCATION: 42572..42591
; OTHER INFORMATION: 20-841.rp complement
; NAME/KEY: primer_bind
; LOCATION: 45328..45347
; OTHER INFORMATION: 20-842.pu
; NAME/KEY: primer_bind
; LOCATION: 45863..45883
; OTHER INFORMATION: 20-842.rp complement
; NAME/KEY: primer_bind
; LOCATION: 76644..76664
; OTHER INFORMATION: 20-853.pu
; NAME/KEY: primer_bind
; LOCATION: 77166..77185
; OTHER INFORMATION: 20-853.rp complement
; NAME/KEY: primer_bind
; LOCATION: 1220..1238
; OTHER INFORMATION: 20-828-311.mis
; NAME/KEY: primer_bind
; LOCATION: 1240..1258
; OTHER INFORMATION: 20-828-311.mis complement
; NAME/KEY: primer_bind
; LOCATION: 12328..12346
; OTHER INFORMATION: 17-42-319.mis
; NAME/KEY: primer_bind
; LOCATION: 12348..12366
; OTHER INFORMATION: 17-42-319.mis complement
; NAME/KEY: primer_bind
; LOCATION: 15242..15260
; OTHER INFORMATION: 17-41-250.mis complement
; NAME/KEY: primer_bind
; LOCATION: 42199..42217
; OTHER INFORMATION: 20-841-149.mis
; NAME/KEY: primer_bind
; LOCATION: 42219..42237
; OTHER INFORMATION: 20-841-149.mis complement
; NAME/KEY: primer_bind
; LOCATION: 45423..45441
; OTHER INFORMATION: 20-842-115.mis
; NAME/KEY: primer_bind
; LOCATION: 45443..45461
; OTHER INFORMATION: 20-842-115.mis complement
; NAME/KEY: primer_bind
; LOCATION: 77039..77057
; OTHER INFORMATION: 20-853-415.mis
; NAME/KEY: primer_bind

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; LOCATION: 77059..77077
; OTHER INFORMATION: 20-853-415.mis complement
; NAME/KEY: misc_binding
; LOCATION: 1227..1251
; OTHER INFORMATION: 20-828-311.probe
; NAME/KEY: misc_binding
; LOCATION: 12335..12359
; OTHER INFORMATION: 17-42-319.probe
; NAME/KEY: misc_binding
; LOCATION: 15229..15253
; OTHER INFORMATION: 17-41-250.probe
; NAME/KEY: misc_binding
; LOCATION: 42206..42230
; OTHER INFORMATION: 20-841-149.probe
; NAME/KEY: misc_binding
; LOCATION: 45430..45454
; OTHER INFORMATION: 20-842-115.probe
; NAME/KEY: misc_binding
; LOCATION: 77046..77070
; OTHER INFORMATION: 20-853-415.probe
; US-09-751-877-1

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Query Match 3.4%; Score 187.6; DB 10; Length 81001;
 Best Local Similarity 71.1%; Pred. No. 1e-38;
 Matches 263; Conservative 104; Indels 3; Gaps 1;

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QY 5084 TCATCATATATTTAAATTTAAATTTCTTAAGAAGAGGCTGGTGCAGTGGCTCAGCCTTTAA 5143
Db 65068 TTACCCCTGATTTTCAAAATAAGGAAATTTGTGGCCAGGTGCAGTGGCTCAGCCTGTAA 65009

QY 5144 TCCAGCACTTTGGGAAGCCAGGAGGAGTCTCTTGAACACAGGAGTTTGACACCAG 5203
Db 65008 TCCAGCACTTTGGGAGGCCAAGGTGGTGGATCACTTGAGTCAAGAGTTGAGACCAG 64949

QY 5204 CTGAGCAACAAAGCAAGCAACCCCATCTCTATAAAAAATTAATAAATTTAGTGGCATGGT 5263
Db 64948 CCTGGCCAAACATGGTGAACCCCGTCTCTATTAAAAATACAAAAATAGCCGGCATGGT 64889

QY 5264 GGCACATGCTGTAGTCCAGCTACTCCAGAGGCTGAG--ATGGATCATCTGAGCCTCA 5320
Db 64888 GGCAGAGCCTGTATCCAGTCTACTCAGGAGGCTGAGGAGGAGAAATCACTTGAACCCG 64829

QY 5321 GGAGTTGAGGCTGCAGTGAGCTGTGACTGCGCCACTGCATCTCCAGTCTGGGACAAACA 5380
Db 64828 GGAGGCGGAGGTTGCAGTGAGCTGAGATTGCGCCACTGCATCTCCAGCTGAGGACAAGA 64769

QY 5381 GCAGACCCCTGCTTTAAAAAAGAAAAAAGAAAAAATTTTTTTCTAAGAAAGCTGTCTTA 5440
Db 64768 GCGAGACTCTGTATCAAAAAAAGAAAAAAGAAAAAATTAATAAAGGAAACTGTGGCTCAG 64709

QY 5441 CAAAGTTGAG 5450
Db 64708 AGAAGTTAAG 64699

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RESULT 15

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US-09-764-872-812/c
; Sequence 812, Application US/09764872
; Publication No. US20030050231A1
; GENERAL INFORMATION:
; APPLICANT: Rosen et al.
; TITLE OF INVENTION: Nucleic Acids, Proteins, and Antibodies
; FILE REFERENCE: PA125
; CURRENT APPLICATION NUMBER: US/09/764,872
; PRIORITY FILING DATE: 2001-01-17
; Prior application data removed - consult PALM or file wrapper
; NUMBER OF SEQ ID NOS: 957
; SOFTWARE: PatentIn Ver. 2.0
; SEQ ID NO 812
; LENGTH: 31718
; TYPE: DNA
; ORGANISM: Homo sapiens
; US-09-764-872-812

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Query Match		3.4%;	Score 187.4;	DB 9;	Length 31718;
Best Local Similarity		75.4%;	Pred. No. 5.9e-39;		
Matches 260;		Conservative	0;	Mismatches 81;	Indels 4; Gaps 2;
Qy	5097	AAATTAAATTTCTAAGNAGAGGCTGGTGCAGTGGCTCACACCTTTAATCCAGCACTTTG	5156		
Db	3273	AAAAGAAACGCTCTCTCGAAGCCGGTGGGTGGCTCACCTGTAAATCCAGCACTTTG	3214		
Qy	5157	GGAAAGCAAAGGAGGAGGACTGCTTGAACACAGGAGTTTGAGACCAGCTGAGCAACAA	5216		
Db	3213	GGAGGCCGAGGAGGCGGATCACCTGAGGTCAGGAGTTGGAGACCAGCTGGCCGACATG	3154		
Qy	5217	GCAAGACCCCATCTCTATAAAACTAAATAATTAAGTTGGGCATGGTGGCAGATCCCTGT	5276		
Db	3153	GCAGAACCCCATCTCTACTATAAAATACAGAAATTAAGCTAGGCATGGTGGCAGATCCCTGT	3094		
Qy	5277	AGTCCCAGCTACTCCAGAGGCTGAG--ATGGATCATCTGAGCCTCAGGAGGTTGAGGCT	5333		
Db	3093	AGCCCAGCTACTTGGGAGGCTGAGGCATAGAATCGTTGAACCCAGGAGGAGGCT	3034		
Qy	5334	GCAGTGAGCTGTGACTGCGCCACTGCATCCAGTCTGGGACAAACAGAGCAAGCCCTGTC	5393		
Db	3033	GCAGTGAGCTGAGACTGTGCCACTGCATCCAGCCTGGG-CAACAGAAAGAGACTCTGTC	2975		
Qy	5394	TTAAAAAAGAAAGAAATTTTTTTCTAAGAGCTGTC	5438		
Db	2974	TCAAAAAAAGAAAGAAATGCTCTCTTTATTCAAGGTACC	2930		

Search completed: June 12, 2003, 04:54:41
Job time : 710 secs